

GTAGCCCTACTCGTAAGG (SEQ ID NO:2) or GAGCAAAGGTATTAACTTACTCCC
(SEQ ID NO:3) or GTTAGCCGTCCCTTCTGG (SEQ ID NO:4).

15. (Amended) A method according to claim 14 wherein said probe is having no more than five, preferably no more than two mismatches with a probe selected of a group composed of probes having a sequence TTATCCCCCTCTGATGGG (SEQ ID NO:5) or AGAGAAGCAAGCTTCTCGTCCG (SEQ ID NO:10) or GCCACTCCTTTTCCGG (SEQ ID NO:7).

19. (Amended) A method according to claim 18 wherein said probe is having no more than five, preferably no more than two mismatches with a probe selected of a group composed of probes having a sequence GCTAATGCAGCGCGATCC (SEQ ID NO:8) or CCGAAGGGGAAGGCTCTA (SEQ ID NO:9) or AGAGAAGCAAGCTTCTCGTCCGTT (SEQ ID NO:10).

21. (Amended) A method according to claim 20 wherein said positive control probe comprising no more than five mismatches with a probe with the sequence GCTGCCTCCGTAGGAGT (SEQ ID NO:11) and/or wherein said negative control probe comprises no more than five mismatches with a probe with the sequence
ACTCCTACGGGGAGGGCACCC (SEQ ID NO:12)